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1635

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1600

RAW SEQUENCE LISTING

DATE: 07/19/2002

PATENT APPLICATION: US/09/780,224B

TIME: 14:49:37

Input Set : A:\208859.ST25.txt

Output Set: N:\CRF3\07192002\I780224B.raw

3 <110> APPLICANT: Roelvink, Petrus W
4 Kovesdi, Imre
5 Wickham, Thomas J
7 <120> TITLE OF INVENTION: ADENOVIRAL CAPSID CONTAINING CHIMERIC PROTEIN IX
9 <130> FILE REFERENCE: 208859
11 <140> CURRENT APPLICATION NUMBER: US 09/780,224B
C--> 12 <141> CURRENT FILING DATE: 2002-07-02
14 <150> PRIOR APPLICATION NUMBER: US 60/181,163
15 <151> PRIOR FILING DATE: 2000-02-09
17 <160> NUMBER OF SEQ ID NOS: 15
19 <170> SOFTWARE: PatentIn Ver. 3.1
21 <210> SEQ ID NO: 1
22 <211> LENGTH: 144
23 <212> TYPE: PRT
24 <213> ORGANISM: Adenovirus
26 <400> SEQUENCE: 1
27 Met Asn Gly Thr Thr Gln Asn Asn Ala Ala Leu Phe Asp Gly Gly Val
28 1 5 10 15
30 Phe Ser Pro Tyr Leu Thr Ser Arg Leu Pro Tyr Trp Ala Gly Val Arg
31 20 25 30
33 Gln Asn Val Val Gly Ser Thr Val Asp Gly Arg Pro Val Ala Pro Ala
34 35 40 45
36 Asn Ser Ser Thr Leu Thr Tyr Ala Thr Ile Gly Pro Ser Pro Leu Asp
37 50 55 60
39 Thr Ala Ala Ala Ala Ala Ser Ala Ala Ala Ser Thr Ala Arg Ser
40 65 70 75 80
42 Met Ala Ala Asp Phe Ser Phe Tyr Asn His Leu Ala Ser Asn Ala Val
43 85 90 95
45 Thr Arg Thr Ala Val Arg Glu Asp Ile Leu Thr Val Met Leu Ala Lys
46 100 105 110
48 Leu Glu Thr Leu Thr Ala Gln Leu Glu Glu Leu Ser Gln Lys Val Glu
49 115 120 125
51 Glu Leu Ala Asp Ala Thr Thr His Thr Pro Ala Gln Pro Val Thr Gln
52 130 135 140
54 <210> SEQ ID NO: 2
55 <211> LENGTH: 125
56 <212> TYPE: PRT
57 <213> ORGANISM: Adenovirus
59 <400> SEQUENCE: 2
60 Met Ala Glu Glu Gly Arg Ile Tyr Val Pro Tyr Val Thr Ala Arg Leu
61 1 5 10 15
63 Pro Lys Trp Ser Gly Ser Val Gln Asp Lys Thr Gly Ser Asn Met Leu
64 20 25 30

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66 Gly Gly Val Val Leu Pro Pro Asn Ser Gln Ala His Arg Thr Glu Thr
67           35                     40                     45
69 Val Gly Thr Glu Ala Thr Arg Asp Asn Leu His Ala Glu Gly Ala Arg
70           50                     55                     60
72 Arg Pro Glu Asp Gln Thr Pro Tyr Met Ile Leu Val Glu Asp Ser Leu
73 65           70                     75                     80
75 Gly Gly Leu Lys Arg Arg Met Asp Leu Leu Glu Glu Ser Asn Gln Gln
76           85                     90                     95
78 Leu Leu Ala Thr Leu Asn Arg Leu Arg Thr Gly Leu Ala Ala Tyr Val
79           100                    105                    110
81 Gln Ala Asn Leu Val Gly Gly Gln Val Asn Pro Phe Val
82           115                    120                    125
84 <210> SEQ ID NO: 3
85 <211> LENGTH: 125
86 <212> TYPE: PRT
87 <213> ORGANISM: Adenovirus
89 <400> SEQUENCE: 3
90 Met Ala Glu Glu Gly Arg Ile Tyr Val Pro Tyr Val Thr Ala Arg Leu
91 1           5                     10                    15
93 Pro Lys Trp Ser Gly Ser Val Gln Asp Lys Thr Gly Ser Asn Met Leu
94           20                     25                     30
96 Gly Gly Val Val Leu Pro Pro Asn Ser Gln Ala His Arg Thr Glu Thr
97           35                     40                     45
99 Val Gly Thr Glu Ala Thr Arg Asp Asn Leu His Ala Glu Gly Ala Arg
100          50                     55                     60
102 Arg Pro Glu Asp Gln Thr Pro Tyr Met Ile Leu Val Glu Asp Ser Leu
103 65           70                     75                     80
105 Gly Gly Leu Lys Arg Arg Met Asp Leu Leu Glu Glu Ser Asn Gln Gln
106           85                     90                     95
108 Leu Leu Ala Thr Leu Asn Arg Leu Arg Thr Gly Leu Ala Ala Tyr Val
109           100                    105                    110
111 Gln Ala Asn Leu Val Gly Gly Gln Val Asn Pro Phe Val
112          115                    120                    125
114 <210> SEQ ID NO: 4
115 <211> LENGTH: 140
116 <212> TYPE: PRT
117 <213> ORGANISM: Adenovirus
119 <400> SEQUENCE: 4
120 Met Ser Ala Asn Ser Phe Asp Gly Ser Ile Val Ser Ser Tyr Leu Thr
121 1           5                     10                    15
123 Thr Arg Met Pro Pro Trp Ala Gly Val Arg Gln Asn Val Met Gly Ser
124           20                     25                     30
126 Ser Ile Asp Gly Arg Pro Val Leu Pro Ala Asn Ser Thr Thr Leu Thr
127           35                     40                     45
129 Tyr Glu Thr Val Ser Gly Thr Pro Leu Glu Thr Ala Ala Ser Ala Ala
130          50                     55                     60
132 Ala Ser Ala Ala Ala Ala Thr Ala Arg Gly Ile Val Thr Asp Phe Ala
133 65           70                     75                     80
135 Phe Leu Ser Pro Leu Ala Ser Ser Ala Ala Ser Arg Ser Ser Ala Arg

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136          85          90          95
138 Asp Asp Lys Leu Thr Ala Leu Leu Ala Gln Leu Asp Ser Leu Thr Arg
139          100          105          110
141 Glu Leu Asn Val Val Ser Gln Gln Leu Leu Asp Leu Arg Gln Gln Val
142          115          120          125
144 Ser Ala Leu Lys Ala Ser Ser Pro Pro Asn Ala Val
145          130          135          140
147 <210> SEQ ID NO: 5
148 <211> LENGTH: 140
149 <212> TYPE: PRT
150 <213> ORGANISM: Adenovirus
152 <400> SEQUENCE: 5
153 Met Ser Thr Asn Ser Phe Asp Gly Ser Ile Val Ser Ser Tyr Leu Thr
154 1          5          10          15
156 Thr Arg Met Pro Pro Trp Ala Gly Val Arg Gln Asn Val Met Gly Ser
157          20          25          30
159 Ser Ile Asp Gly Arg Pro Val Leu Pro Ala Asn Ser Thr Thr Leu Thr
160          35          40          45
162 Tyr Glu Thr Val Ser Gly Thr Pro Leu Glu Thr Ala Ala Ser Ala Ala
163          50          55          60
165 Ala Ser Ala Ala Ala Ala Thr Ala Arg Gly Ile Val Thr Asp Phe Ala
166 65          70          75          80
168 Phe Leu Ser Pro Leu Ala Ser Ser Ala Ala Ser Arg Ser Ser Ala Arg
169          85          90          95
171 Asp Asp Lys Leu Thr Ala Leu Leu Ala Gln Leu Asp Ser Leu Thr Arg
172          100          105          110
174 Glu Leu Asn Val Val Ser Gln Gln Leu Leu Asp Leu Arg Gln Gln Val
175          115          120          125
177 Ser Ala Leu Lys Ala Ser Ser Pro Pro Asn Ala Val
178          130          135          140
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181 <211> LENGTH: 132
182 <212> TYPE: PRT
183 <213> ORGANISM: Adenovirus
185 <400> SEQUENCE: 6
186 Met Ser Gly Phe Thr Glu Gly Asn Ala Val Ser Phe Glu Gly Gly Val
187 1          5          10          15
189 Phe Ser Pro Tyr Leu Thr Thr Arg Leu Pro Ser Trp Ala Gly Val Arg
190          20          25          30
192 Gln Asn Val Val Gly Ser Asn Val Asp Gly Arg Pro Val Ala Pro Ala
193          35          40          45
195 Asn Ser Thr Thr Leu Thr Tyr Ala Thr Ile Gly Ser Ser Val Asp Thr
196          50          55          60
198 Ala Ala Ala Ala Ala Ala Ser Ala Ala Ala Ser Thr Ala Arg Gly Met
199 65          70          75          80
201 Ala Ala Asp Phe Gly Leu Tyr Asn Gln Leu Ala Ala Ser Arg Leu Arg
202          85          90          95
204 Glu Glu Asp Ala Leu Ser Val Val Leu Thr Arg Leu Glu Glu Leu Ser
205          100          105          110

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207 Gln Gln Leu Gln Asp Met Ser Ala Lys Met Ala Leu Leu Asn Pro Pro
208      115      120      125
210 Ala Asn Thr Ser
211      130
213 <210> SEQ ID NO: 7
214 <211> LENGTH: 133
215 <212> TYPE: PRT
216 <213> ORGANISM: Adenovirus
218 <400> SEQUENCE: 7
219 Met Ser Gly Ser Met Glu Gly Asn Ala Val Ser Phe Lys Gly Gly Val
220 1      5      10      15
222 Phe Ser Pro Tyr Leu Thr Thr Arg Leu Pro Ala Trp Ala Gly Val Arg
223      20      25      30
225 Gln Asn Val Met Gly Ser Asn Val Asp Gly Arg Pro Val Ala Pro Ala
226      35      40      45
228 Asn Ser Ala Thr Leu Thr Tyr Ala Thr Val Gly Ser Ser Val Asp Thr
229      50      55      60
231 Ala Ala Ala Ala Ala Ala Ser Ala Ala Ala Ser Thr Ala Arg Gly Met
232 65      70      75      80
234 Ala Ala Asp Phe Gly Leu Tyr Asn Gln Leu Ala Ala Ser Arg Ser Leu
235      85      90      95
237 Arg Glu Glu Asp Ala Leu Ser Val Val Leu Thr Arg Met Glu Glu Leu
238      100      105      110
240 Ser Gln Gln Leu Gln Asp Leu Phe Ala Lys Val Ala Leu Leu Asn Pro
241      115      120      125
243 Pro Ala Asn Ala Ser
244      130
246 <210> SEQ ID NO: 8
247 <211> LENGTH: 130
248 <212> TYPE: PRT
249 <213> ORGANISM: Adenovirus
251 <220> FEATURE:
252 <221> NAME/KEY: misc_feature
253 <222> LOCATION: (2)..(5)
254 <223> OTHER INFORMATION: "Xaa" may be any amino acid
256 <220> FEATURE:
257 <221> NAME/KEY: misc_feature
258 <222> LOCATION: (7)..(7)
259 <223> OTHER INFORMATION: "Xaa" may be any amino acid
261 <220> FEATURE:
262 <221> NAME/KEY: misc_feature
263 <222> LOCATION: (9)..(9)
264 <223> OTHER INFORMATION: "Xaa" may be any amino acid
266 <220> FEATURE:
267 <221> NAME/KEY: misc_feature
268 <222> LOCATION: (11)..(11)
269 <223> OTHER INFORMATION: "Xaa" may be any amino acid
271 <220> FEATURE:
272 <221> NAME/KEY: misc_feature

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Input Set : A:\208859.ST25.txt

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273 <222> LOCATION: (21)..(21)
274 <223> OTHER INFORMATION: "Xaa" may be any amino acid
276 <220> FEATURE:
277 <221> NAME/KEY: misc_feature
278 <222> LOCATION: (30)..(30)
279 <223> OTHER INFORMATION: "Xaa" may be any amino acid
281 <220> FEATURE:
282 <221> NAME/KEY: misc_feature
283 <222> LOCATION: (34)..(34)
284 <223> OTHER INFORMATION: "Xaa" may be any amino acid
286 <220> FEATURE:
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288 <222> LOCATION: (45)..(45)
289 <223> OTHER INFORMATION: "Xaa" may be any amino acid
291 <220> FEATURE:
292 <221> NAME/KEY: misc_feature
293 <222> LOCATION: (54)..(57)
294 <223> OTHER INFORMATION: "Xaa" may be any amino acid
296 <220> FEATURE:
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298 <222> LOCATION: (69)..(69)
299 <223> OTHER INFORMATION: "Xaa" may be any amino acid
301 <220> FEATURE:
302 <221> NAME/KEY: misc_feature
303 <222> LOCATION: (74)..(76)
304 <223> OTHER INFORMATION: "Xaa" may be any amino acid
306 <220> FEATURE:
307 <221> NAME/KEY: misc_feature
308 <222> LOCATION: (79)..(83)
309 <223> OTHER INFORMATION: "Xaa" may be any amino acid
311 <220> FEATURE:
312 <221> NAME/KEY: misc_feature
313 <222> LOCATION: (86)..(86)
314 <223> OTHER INFORMATION: "Xaa" may be any amino acid
316 <220> FEATURE:
317 <221> NAME/KEY: misc_feature
318 <222> LOCATION: (88)..(90)
319 <223> OTHER INFORMATION: "Xaa" may be any amino acid
321 <220> FEATURE:
322 <221> NAME/KEY: misc_feature
323 <222> LOCATION: (92)..(95)
324 <223> OTHER INFORMATION: "Xaa" may be any amino acid
326 <220> FEATURE:
327 <221> NAME/KEY: misc_feature
328 <222> LOCATION: (98)..(98)
329 <223> OTHER INFORMATION: "Xaa" may be any amino acid
331 <220> FEATURE:
332 <221> NAME/KEY: misc_feature
333 <222> LOCATION: (100)..(101)

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/780,224B

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Input Set : A:\208859.ST25.txt
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:8; Xaa Pos. 3,4,5,7,9,11,21,30,34,45,54,55,56,57,69,74,75,76,79,80,81

Seq#:8; Xaa Pos. 82,83,86,88,89,90,92,93,94,95,98,100,101,105,107,108,110

Seq#:8; Xaa Pos. 111,112,114,115,116,119,120,122,123,124,125,127,129

VERIFICATION SUMMARY

DATE: 07/19/2002

PATENT APPLICATION: US/09/780,224B

TIME: 14:49:38

Input Set : A:\208859.ST25.txt

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L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:377 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
L:380 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:16
L:383 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:32
L:386 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:48
L:389 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:64
L:392 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:80
L:395 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:96
L:398 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:112
L:401 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:128